

*AK:shw*

CRF Errors Corrected by the STIC Systems Branch

Serial Number: 09/270,910

ENTERED

CRF Processing Date: 9/5/2000  
Edited by: AK  
Verified by: AK (STIC staff)

RECEIVED

SEP 08 2000

TECH CENTER 1600/2900

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: \_\_\_\_\_
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other \_\_\_\_\_
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: \_\_\_\_\_
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: \_\_\_\_\_
- ☒ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: 40
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: \_\_\_\_\_
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: \_\_\_\_\_
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as \_\_\_\_\_
- ☐ Inserted mandatory headings, specifically: \_\_\_\_\_
- ☐ Corrected an obvious error in the response, specifically: \_\_\_\_\_
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: \_\_\_\_\_
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: \_\_\_\_\_
- ☐ Other: \_\_\_\_\_

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

M. DiBrino

1644

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/270,910  
DATE: 09/05/2000  
TIME: 11:38:38  
Input Set : A:\Seqlist.txt  
Output Set: N:\CRF3\09052000\I270910.raw

Does Not Comply  
Corrected Diskette Needed

4 <110> APPLICANT: Ipsen, Hans Henrick  
5 Spangfort, Michael Dho  
6 Larsen Jorgen Nedergaard  
8 <120> TITLE OF INVENTION: NOVEL RECOMBINANT ALLERGENS  
11 <130> FILE REFERENCE: 4305/1E144 US1  
13 <140> CURRENT APPLICATION NUMBER: 09/270,910  
14 <141> CURRENT FILING DATE: 1999-03-16  
16 <150> PRIOR APPLICATION NUMBER: 60/078,371  
17 <151> PRIOR FILING DATE: 1998-03-18  
19 <160> NUMBER OF SEQ ID NOS: 40  
21 <170> SOFTWARE: FastSEQ for Windows Version 3.0

ERRORED SEQUENCES

W--> 501 <210> SEQ ID NO: 40  
502 <211> LENGTH: 38  
503 <212> TYPE: DNA  
504 <213> ORGANISM: Artificial Sequence  
506 <220> FEATURE:  
507 <223> OTHER INFORMATION: primer  
509 <400> SEQUENCE: 40  
E--> 510 ccgctcgaga aaagaacaa ttattgtaaa ataaaaatg

38 ←

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/270,910

DATE: 09/05/2000  
TIME: 11:38:39

Input Set : A:\Seqlist.txt

Output Set: N:\CRF3\09052000\I270910.raw

L:49 M:258 W: Mandatory Feature missing, <220> FEATURE:  
L:501 M:283 W: Missing Blank Line separator, <210> field identifier  
L:510 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:38 SEQ:40